

Glycopattern analysis and structure of the egg extra-cellular matrix in the Apennine yellow-bellied toad, *Bombina pachypus* (Anura: Bombinatoridae)

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Abstract: We studied the glycopatterns and ultrastructure of the extra-cellular matrix (ECM) of the egg of the Apennine yellow-bellied toad *Bombina pachypus*, by light and electron microscopy in order to determine structure, chemical composition and function. Histochemical techniques in light microscopy included PAS and Alcian Blue pH 2.5 and 1.0, performed also after β -elimination. Lectin-binding was tested with nine lectins (AAA, ConA, DBA, HPA, LTA, PNA, SBA, UEA-I, WGA). An inner fertilization envelope (FE) and five jelly layers (J_1 – J_5) were observed, differing in histochemical staining, lectin binding and ultrastructure. Most glycans were O-linked, with many glucosylated and fucosylated residues. The fertilization envelope presented a perivitelline space and a fertilization layer, with mostly neutral glycans. The jelly layers consisted of fibers and granules, whose number and orientation differed between layers. Fibers were densely packed in J_1 and J_4 layers, whereas a looser arrangement was observed in the other layers. Jelly-layer glycans were mostly acidic and particularly abundant in the J_1 and J_4 layers. In the J_1 , J_2 and J_5 layers, neutral, N-linked glycans were also observed. Mannosylated and/or glucosylated as well as galactosyl/galactosaminylated residues were more abundant in the outer layers. Many microorganisms were observed in the J_5 layer. We believe that, apart from their functions in the fertilization process, acidic and fucosylated glycans could act as a barrier against pathogen penetration. (*Folia Histochemica et Cytobiologica* 2011; Vol. 49, No. 2, pp. 306–316)

Key words: amphibia, *Bombina pachypus*, egg jelly, lectin, electron microscopy

Introduction

Amphibian eggs are surrounded by an extra-cellular matrix (ECM) consisting of an inner vitelline envelope and an outer jelly coat, made up of layers whose number and composition are species-specific

[1–3]. The jelly coat layers consist mainly of glycoproteins and glycosaminoglycans [3–5], and their carbohydrate-chain composition has been investigated in a number of species [6–9]. In general, the glycans are O-linked to the core proteins and several sequences are unique to a given species [5–11]. In most studies, the whole jelly coat is taken for biochemical analysis, without distinguishing between the layers or the sublayers. Histochemical studies have demonstrated that the distribution of neutral and acidic carbohydrates varies among the layers in several species [1, 4, 12]. Lectin-binding experiments on jelly coat layers have revealed heterogeneity in the distribution of binding sites among

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and within layers [5, 13, 14]. Electron-microscopy studies have revealed that each layer presents a unique fiber and glycoprotein composition, and that the jelly coats are complex structures in which globular glycoproteins are bound to a fibrous glycoprotein superstructure [5, 15].

Jelly coat layers have been found to be involved in a number of functions, such as interactions with spermatozoa in fertilization [16, 17], mechanical support, spacing and substrate attachment for the egg clusters [18], hosting for oxygen-supplying, symbiotic algae [19], and protection against predators, pathogens, environmental stressors, and contaminants [3, 20, 21]. Amphibians with a more complex jelly coat covering show a higher resistance to water molds [22]. The penetration of contaminants is significantly reduced and/or slowed by jelly coat layers [21, 23], even if this is not always observed [24, 25]. The resistance of egg ECM to environmental stresses can even vary geographically within a single species [26]. Several papers have attempted to link the presence of particular molecules in the ECM to given functions [e.g. 5, 8, 10, 11, 13, 16, 27]. A relatively small number of species has been investigated taking into account the huge diversity of ECM organization among living amphibians [1, 3, 18], and the role of several jelly molecules is still poorly understood [27].

Knowledge about the structure and composition of the egg jelly coat is fundamental to understanding both the fertilization processes and the mechanisms by which contaminants, e.g. pesticides and pathogens, e.g. Saprolegniaceae, affect egg survival and the consequent reproductive success of amphibian species, the prevalence of many of which is declining around the world [28, 29].

With all this in mind, in the present paper we studied, using different staining and microscopic observation techniques, the composition and organization of ECM in the Apennine yellow-bellied toad, *Bombina pachypus*.

Our goals were as follows:

1. To link the structure and glycopolymers composition of such a complex structure;
2. To give some functional interpretations of the findings in the light of the available literature;
3. To obtain preliminary data in understanding the role of ECM in the protection of the egg, and its implications for future studies about the effects of contaminants on the survival of eggs in this endemic, but declining, species [30];
4. To contribute to the understanding of ECM diversity among anurans.

Material and methods

Egg collection. Eggs of the Apennine yellow-bellied toad, *Bombina pachypus*, were collected from fields in the Gravine of Laterza (Taranto, Apulia, Italy) in April 2008 soon after their deposition. No more than one pair of eggs was taken from each clutch, usually consisting of 15–20 eggs, to avoid hampering reproductive success. This egg collection was authorized by Italy's Ministero dell'Ambiente.

Light microscopy. The eggs were embedded in a Technovit 8100 kit (EMS, Hatfield, PA, USA). The eggs were fixed in a 4% paraformaldehyde solution in 0.1 M phosphate-buffered-saline (PBS) pH 7.4 at 4°C for three hours. After several rinses in PBS, pieces were incubated overnight, at 4°C, in PBS with 6.8% added sucrose, and then dehydrated with increasing acetone, also at 4°C. Infiltration was performed by incubating the specimens in a Technovit 8100 monomer for six hours at 4°C with gentle stirring. Finally, the eggs were embedded with an ice-cold solution of 15:1 infiltrating solution. Polymerization was carried out on an ice bed for three hours. Semi-thin sections (2 µm thick) were cut with glass knives using an LKB Ultratome and mounted on microscope slides, coated with polylysine (Sigma, St. Louis, MO, USA). Semi-thin sections were incubated for five minutes at 37°C in 0.01% trypsin (Sigma) and 0.1% CaCl₂ in PBS, pH 7.8 before staining.

Histochemistry. The sections were stained with periodic acid-Schiff (PAS) hemallum [31] for general carbohydrate staining, and Alcian-Blue (AB) at pH 2.5 or pH 1.0 to detect acidic carbohydrates. All the cited reagents were from Sigma. PAS and Alcian stainings were also combined to differentiate between neutral and acidic carbohydrates in the same section [32]. PAS-AB pH 2.5 were also performed after β-elimination, a method that removes the O-linked oligosaccharides from glycoproteins [33]; prior to staining, sections were incubated with 0.2M KOH in dimethylsulphoxide — H₂O — ethanol (50:40:10) for one hour at 45°C, followed by neutralisation with 10 mM HCl and washing in PBS pH 7.4.

Lectin histochemistry. The binding of nine lectins (all from Sigma except for AAA from Vector Laboratories, Burlingame, CA, USA) was assessed to determine the nature and distribution of glycosidic residues in the egg layers. Lectins were labelled with horseradish peroxidase (HRP), fluoresceine isothiocyanate (FITC) or phosphatase. The lectins, their concentrations, and their sugar specificities are summarized in Table 1. References for the lectins are given in References 34–42.

For the binding with FITC-conjugated lectins (ConA, DBA, HPA, PNA, LTA, WGA, UEA-I), the sections were incubated for 30 minutes in blocking buffer, i.e. 1% normal

Table 1. Characteristics of the lectins utilized

Lectin	Source and reference numbers	Binding specificity	Lectin concentration [mg/ml]	Inhibitory sugar
Con A	<i>Canavalia ensiformis</i> [34]	D-mannose, D-Glucose	0.005	0.1 M M α M
WGA	<i>Triticum vulgaris</i> [35]	(GlcNAc β 1,4) _n	0.02	0.01 M TACT
SBA	<i>Glycine max</i> [36]	GalNAc	0.02	0.2 M GalNAc
HPA	<i>Helix pomatia</i> [37]	GalNAc	0.02	0.2 M GalNAc
DBA	<i>Dolichos biflorus</i> [38]	α -GalNAc	0.02	0.2 M GalNAc
PNA	<i>Arachis hypogaea</i> [39]	Gal β 1,3GalNAc	0.01	0.2 M Gal
AAA	<i>Aleuria aurantia</i> [40]	Fuc α (1,6)GlcNAc- β NA _{sn} Fuc α (1,3), Fuc α (1,4)	0.01	0.2 M L-Fuc
UEA-I	<i>Ulex europaeus</i> [41]	Fuc α (1,2)	0.01	0.2 M L-Fuc
LTA	<i>Tetragonolobus purpureus</i> [42]	L-Fuc α 1,6GlcNAc and L-Fuc α 1,2Gal β 1,4[L-Fuc1,3] GlcNAc β 1,6R	0.02	0.2 M L-Fuc

Fuc — fucose; Gal — galactose; GalNAc — N-acetylgalactosamine; GlcNAc — N-acetylglucosamine; M α M — methyl- α -mannopyranoside; TACT — N,N',N''-triacetylchitotriose

goat serum in 0.1 M Tris-buffered saline pH 7.4 (TBS) and then incubated for one hour at room temperature with the FITC-lectin solution in TBS. Sections were subsequently rinsed in the same buffer and mounted in 70% glycerin in TBS. For the binding with HRP-conjugated SBA lectin, the sections were exposed to 3% hydrogen peroxide for ten minutes to inhibit endogenous peroxidase activity, and then incubated for 60 minutes at room temperature with HRP-lectin in TBS. HRP activity was then visualised with 0.005% 3–3'-diaminobenzidine (DAB; Sigma)-0.01% hydrogen peroxide in 0.05 M TBS [43] for ten minutes in the dark at room temperature. Finally, the sections were dehydrated through a graded ethanol series, cleared in Histolemon (Carlo Erba, Rodano, Milan, Italy), and mounted in DPX (Fluka BioChemika, Steinheim, Germany).

For the binding with AAA phosphate-conjugated lectin, the sections were incubated for one hour at room temperature with the lectin solution in TBS. Sections were subsequently rinsed in the same buffer and incubated in the substrate working solution (BCIP/NBT alkaline phosphatase substrate Kit IV from Vector Laboratories, Burlingame, CA, USA) for 15 minutes at room temperature. The endogenous alkaline phosphatase activity was inhibited by adding levamisole to the working solution. After washing in 0.1 M TBS pH 9.5 for five minutes, the sections were counterstained with methyl green (Sigma) dehydrated, cleared and mounted following the HRP-lectin protocol.

Two different controls for lectin labeling were used: 1) substitution of the respective lectin with TBS alone; and 2) incubation in the lectin with the addition of the appropriate inhibitory sugar (concentrations are set out in Table 1). Positive controls were included from different regions of the digestive system from two amphibians, *Bufo balearicus* (formerly known as *Bufo viridis*) and *Triturus carnifex*, whose mucins are known to bind to the tested lectins [44, 45].

Transmission electron microscopy. The eggs were fixed in 4% glutaraldehyde and processed for embedding in Epoxy Resin-Araldite (M) CY212 (TAAB, Aldermaston, UK) as previously reported [46]. Semi-thin sections 2 μ m thick were stained with Toluidine blue-PAS (PAS-TB). Ultra-thin sections were mounted on formwar-coated nickel grids and stained routinely with uranyl acetate and lead citrate [47]. Images were captured using a Nikon Eclipse 600 photomicroscope equipped with a Nikon DMX 1200 camera (Nikon Instruments SpA, Calenzano, Florence, Italy).

Each experiment was repeated twice on specimens taken from three different eggs, giving a total of six repetitions. Staining/labeling in each experiment was assessed by at least two independent observers and scored as positive (+), moderately positive (– +), or negative (–) according to their intensity.

Results

Freshly-collected eggs of *B. pachypus* were surrounded by an ECM about 3–4 mm thick. Embedding reduced the ECM thickness because of dehydration. Staining techniques revealed an inner fertilization envelope (FE), and a jelly coat subdivided into five main layers, called J₁–J₅, with J₁ being the innermost.

Histochemistry

Table 2 summarizes the histochemical staining patterns observed. Figures 1A and 1B show the ECM stained by PAS combined with AB pH 2.5 and pH 1.0, respectively. The FE is weakly PAS-positive and negative with AB at both pH 2.5 and 1.0. PAS-positivity of FE increases towards the J₁ layer, where a more intensely stained sublayer is observed. The J₁ layer

Table 2. Histochemical stainings of the egg extra-cellular matrix of *Bombina pachypus*

Layer	Staining			
	PAS	AB pH 2.5	AB pH 1.0	β -elimination PAS AB pH 2.5
FE	+ -/+ + ¹	--	--	--
J ₁	++	++	++	Red
J ₂	+-	+-	+-	--
J ₃	--	++	++	--
J ₄	++	+ -/+ + ²	+-	Red
J ₅	++	+-	+-	Red

Layers: FE — fertilization envelope; J₁–J₅ — jelly coat layers; ¹PAS-positivity increases in a sublayer at the boundary with J₁;

²Alcianophily increases in a sublayer at the boundary with J₅

Table 3. Lectin binding of the egg extra-cellular matrix of *Bombina pachypus*

Layer	Lectin								
	Con A	WGA	SBA	HPA	DBA	PNA	AAA	UEA-I	LTA
FE	--	+-	--	+-	--	++	+-	--	--
J ₁	+-	++	+-	--	--	--	++	--	+-
J ₂	--	++	--	--	--	--	+-	--	--
J ₃	+-	++	--	++	--	--	+-	--	--
J ₄	++	++	--	++	+-	++	++	--	+-
J ₅	++	++	--	--	+-	++	+-	+-	+-

Layers: FE — fertilization envelope; J₁–J₅ — jelly coat layers

stains deep blue with combined PAS-AB pH 2.5, whereas it stains violet with PAS-AB pH 1.0, suggesting that PAS-positivity is masked by intense alcianophily with PAS-AB pH 2.5. The J₂ layer is made up of a network of fibers and granules positive to PAS and AB stainings. The J₃ layer stains mostly blue at both PAS-AB pH 2.5 and PAS-AB pH 1.0. The J₄ layer is positive to PAS and weakly positive to the AB stainings, with a stronger alcianophilic sublayer at the transition zone with the J₅ layer. The latter has a ‘foamy’ appearance and hosts several microorganisms, with fibers and granules positive to PAS and/or to AB stainings. After β -elimination, only the J₁, J₄ and J₅ layers keep their PAS-positivity, but not their alcianophily, whereas the other layers are not stained or weakly stained (Figure 1C).

Lectin histochemistry

Table 3 sets out the lectin-binding patterns observed. Figures 1D–L show the binding patterns of each lectin. Of the nine lectins tested, AAA (Figure 1D) and WGA (Figure 1H) bind to both the FE and all the jelly layers, whereas the others show selectivity towards one or more layers. Each layer binds to at least

three different lectins. In general, the J₁, J₄ and J₅ layers bind to a higher number of lectins than J₂ and J₃, even if it is not always easy to state the positivity of layers with a loose texture, like J₂, J₃, and J₅. PNA (Figure 1J) and DBA (Figure 1K) bind mostly to the outer jelly layers. UEA-I binds weakly only to the J₅ layer (Figure 1E). No labeling was observed in control sections exposed to DAB-H₂O₂ medium after substitution of TBS for lectin or incubated with the corresponding hapten sugars (Figure 1G).

Transmission electron microscopy

The FE shows two areas, the innermost being less electron-dense than the outer (Figure 2A). In the inner area, sparse granules are observed, whereas in the outer there is a loose mesh of fibers with no clear orientation. In the jelly envelope, each of the five layers observed in light microscopy presents a peculiar structure. J₁ is made up of densely-packed fibers running parallel to the egg’s surface (Figures 2A, B). Fibers with the same orientation of J₁ are also seen in the J₂ layer, but they are more loosely arranged and present a series of granules (Figure 2B). In the J₃ layer, the number of fibers decreases and their orientation is less

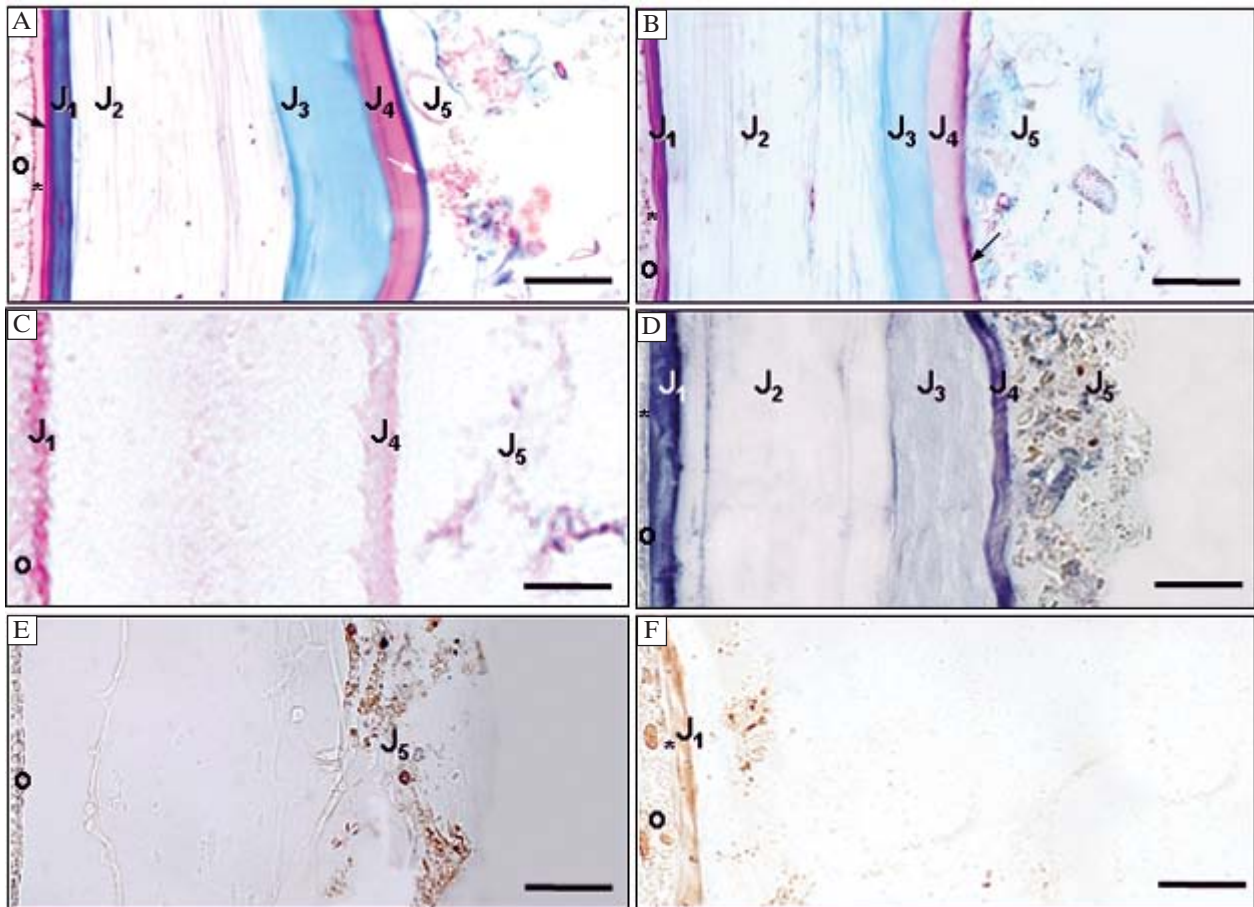


Figure 1. Egg extra-cellular matrix of *Bombina pachypus*. **A.** PAS-Alcian Blue pH 2.5. Layers J₁, J₃ and a sublayer between J₄ and J₅ (white arrow) are intensely alcianophilic. An intensely PAS-positive area is seen in the fertilization envelope (*) towards J₁ (black arrow). J₁–J₅ — jelly coat layers 1 to 5; o, egg. **B.** PAS-Alcian Blue pH 1.0. J₃ is still markedly alcianophilic. The sublayer between J₄ and J₅ is still evident. J₁–J₅ — jelly coat layers 1 to 5; o, egg; * — fertilization envelope. **C.** PAS-Alcian Blue pH 2.5 after b-elimination. PAS-positivity persists in J₁, J₄ and J₅ layers, whereas alcianophily is suppressed. J₁–J₅ — jelly coat layers 1 to 5; o, egg. **D.** Binding with AAA lectin (phosphatase-conjugated). J₁ and J₄ are intensely stained. Several microorganisms can be seen in J₅ layer. J₁–J₅ — jelly coat layers 1 to 5; o, egg; * — fertilization envelope. **E.** Binding with UEA-I lectin (peroxidase-conjugated). Moderate binding can be seen only in J₅ layer. J₅ — jelly coat layer 5; o, egg. **F.** Binding with SBA lectin (peroxidase-conjugated). Moderate binding can be seen only in the J₁ layer. J₁ — jelly coat layer 1; o, egg; * — fertilization envelope

clear, whereas the granular component is more abundant (Figures 2C, D). Densely-packed fibers are again visible in the J₄ layer, the organization of which looks similar to the J₁ layer (Figures 2 D, E). The J₅ layer is characterized by several granules and a loose network of fibers, hosting several microorganisms (Figures 2 E, F). It was not possible to observe the two sublayers revealed by histochemical techniques at the FE/J₁ and J₄/J₅ boundaries, respectively: in these areas, the layers trespass one into another rather abruptly, with no particular structural organizations suggesting the existence of the cited sublayers (Figures 2A, E).

Discussion

By combining histochemical and lectin-histochemical techniques in light microscopy and electron mi-

croscopy, we showed that the extra-cellular matrix of the egg of *B. pachypus* is a complex structure with an envelope and five jelly layers, each characterized by a specific structure and carbohydrate composition.

Lectin-binding patterns do not suggest the presence of a monosaccharide in a glycan, but rather of an oligomer in which a specific monosaccharide is probably present. In any case, lectin-histochemistry is very useful in comparing structures and detecting variations between different layers of the egg envelope [5, 13, 14].

The fertilization envelope, the innermost layer of the ECM, is made up of two zones with different textures, the inner comprising sparse granules and the outer comprising a loose mesh of fibers. The inner and outer zones should correspond to the perivitelline space and the fertilization layer, respectively, observed in *Xenopus laevis* [48]. Glycoconjugates in the FE are

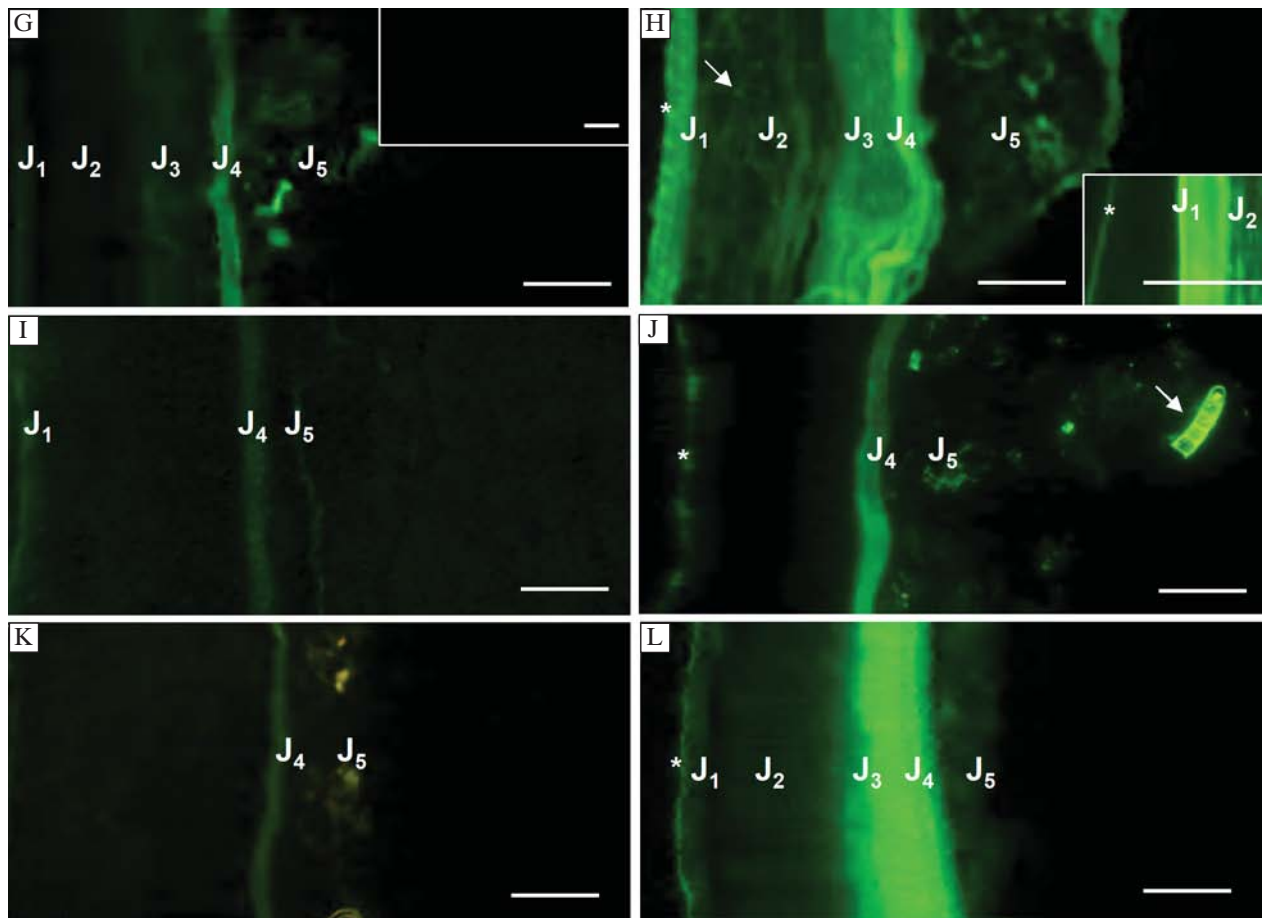


Figure 1 — continued. **G.** Binding with ConA lectin (FITC-conjugated). Binding is seen mostly in J_4 and J_5 layers. Insert: negative control. J_1 – J_5 — jelly coat layers 1 to 5. **H.** Binding with WGA lectin (FITC-conjugated). The lectin binds to all layers. In the J_2 layer lectin binds to both fibers and granules (white arrow). Insert: section with partial detachment of J_1 to show lectin binding to the fertilization envelope (*). J_1 – J_5 — jelly coat layers 1 to 5; * — fertilization envelope. **I.** Binding with LTA lectin (FITC-conjugated). The lectin binds to J_1 , J_4 and J_5 layers. J_1 – J_5 — jelly coat layers 1 to 5; * — fertilization envelope. **J.** Binding with PNA lectin (FITC-conjugated). Binding is mainly observed to the fertilization envelope, J_4 and J_5 . White arrow indicates a fragment of a filamentous green alga. J_1 – J_5 — jelly coat layers 1 to 5; * — fertilization envelope. **K.** Binding with DBA lectin (FITC-conjugated). J_4 and J_5 layers bind to the lectin. J_4 — jelly coat layer 4; J_5 — jelly coat layer 5. **L.** Binding with HPA lectin (FITC-conjugated). Main binding is seen to the fertilization envelope, J_3 and J_4 layers. J_1 – J_5 — jelly coat layers 1 to 5; * — fertilization envelope. Scale bar = 50 μ m

mostly neutral and O-linked, presenting galactosyl/galactosaminylated, glucosaminylated and fucosylated residues. Towards the J_1 layer, a zone that stains more intensely can be seen; boundary zones like these should be referred to as ‘membranes’ [3] and it is unclear whether they are actual, distinct structures or simply optical representations of transition between two layers that have different densities. The lack of differences in lectin binding or structure of this membrane in respect to the FE or the J_1 layer does not support the existence of a true layer between them.

The J_1 layer of *B. pachypus* consists of a number of densely-packed fibers running parallel to the egg’s surface. It is rich in acidic oligosaccharidic residues, as demonstrated by its intense staining with AB pH 2.5. Reduction of alcianophilicity with AB pH 1.0 suggests that the acidic residues are mostly sialylated,

whereas its suppression after β -elimination indicates that such residues are O-linked. By contrast, the persistence of PAS-positivity after β -elimination indicates the presence of neutral, N-linked glycans. Besides, lectin-binding experiments indicate the presence of mannosylated and/or glucosylated, galactosaminylated, and fucosylated residues. WGA-binding could be explained by either the presence of N-acetylglucosamine or Neu5Ac sialic acid.

The J_2 layer has a loose structure, with fibers arranged in a parallel orientation similar to J_1 and granules. Neutral and acidic, mostly O-linked glycans are present, with N-acetylglucosamine and fucose residues in the oligosaccharidic chains.

The J_3 layer presents a fibrous structure like the previous layers, but the fibers are more spaced, with a less clear orientation and an increased number of

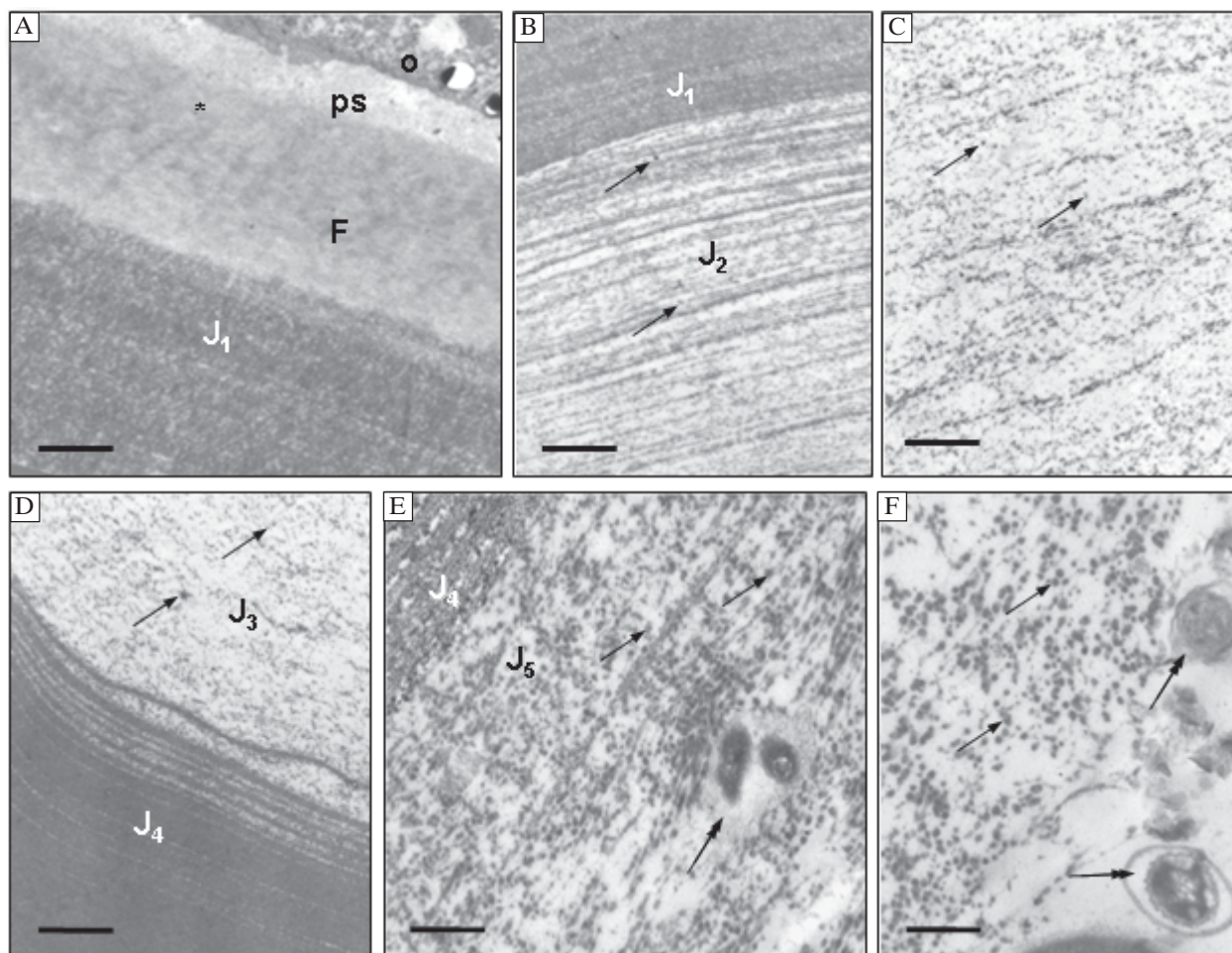


Figure 2. Ultrastructure of egg extra-cellular matrix of *Bombina pachypus* in transmitted electron microscopy. **A.** Fertilization envelope (*) showing the perivitelline space (ps) next to the egg (o) and the fertilization layer (F), and J₁ layer with densely fibrous packing. Scale bar = 1.4 μ m. **B.** J₁ (upper) and J₂ (lower) layers, with the latter having a looser arrangement of fibers and granules (arrows). Scale bar = 1.8 μ m. **C.** J₃ layer with granules (arrows) along fibers. Scale bar = 1.8 μ m. **D.** J₃ (upper right) and J₄ (lower left) layers, the fibers of the latter being densely packed. Arrows indicate granules in the J₃ layer. Scale bar = 1.8 μ m. **E.** Transition between J₄ and J₅ layers, with a reduced number of fibers and an increased number of granules (simple arrows) in the latter. The double arrow indicates a microorganism, possibly a blue alga. Scale bar = 0.7 μ m. **F.** Detail of J₅ layer with granules (simple arrows) and two microorganisms (double arrows). Scale bar = 0.45 μ m

granules. It stains blue with both AB pH 2.5 and AB pH 1.0, and is PAS-negative, whereas β -elimination suppresses alcianophilicity. These results suggest that the fibers contain proteoglycans with carboxylated and sulfated glycosaminoglycans O-linked to the core protein, since both proteoglycans and glycosaminoglycans are scarcely, or not at all, PAS-positive [49]. Residues of mannose and/or glucose, N-acetylglucosamine, N-acetylgalactosamine and fucose are probably present in the oligosaccharidic chains.

Similar to J₁, the J₄ layer has densely-packed fibers. These are PAS-positive and weakly alcianophilic with AB pH 2.5. The suppression of alcianophilicity with AB pH 1.0 and β -elimination indicate the presence of carboxylated, O-linked oligosaccharidic chains.

Several lectins bind to this layer, suggesting the presence of mannosylated, glucosyl/glucosaminylated, galactosyl/galactosaminylated and fucosylated residues. Similar to that observed at the FE/J₁ boundary, a 'membrane' can be seen between J₄ and J₅ intensely stained with PAS-AB pH 2.5 and PAS-AB pH 1.0, but the lack of distinctive ultrastructural and lectin-binding features in respect of the bordering layers suggests that this membrane is merely an optical transition between the layers.

The J₅ layer appears foamy in light microscopy, being made up of a very loose network of fibers and granules. The J₅ layer is involved in the spacing and adhesion of the eggs among them and to the substrate. Neutral and acidic glycans, both O- and N-linked, are

present, with residues of mannose and/or glucose, galactose, N-acetyl-glucosamine, N-acetyl-galactosamine, and fucose. Several microorganisms found at this level can sometimes confuse the interpretation of lectin-binding to the J₅ layer because they are coated with autogenous mucins.

In the jelly envelopes of the species *Bombina bombina* and *B. variegata* (*B. pachypus* is sometimes included in the latter species [50]), 28 O-linked carbohydrate chains have been identified [7]. The species have very similar glycans, sharing the sequences GlcNAc(β1-3)[Fuc(α1-4)]GlcNAc(β1-6) and GlcNAc(α1-4)Gal(β1-4)Gal(β1-3), and fucosylated residues are linked mostly in (1,4) to proximal N-acetylglucosamine or in (1,2) to galactose [7]. The presence of similar residues in *B. pachypus* can be inferred from our lectin-binding experiments, but we cannot compare the results for each single layer with those from *B. bombina* and *B. variegata*, because their jelly envelopes were analyzed as a whole, without distinction between layers [7].

The fertilization layer in the FE of *B. pachypus* differs from that of *X. laevis* in lacking both a clear parallel orientation of the fibers and a more electron-dense F_c layer [48]. The latter is apparently also missing from the FE of other species [4, 51]. It is unclear whether this layer is really missing in *B. pachypus* or was not revealed by our techniques. The filamentous structures in the FE are interpreted as aggregates of ZP glycoproteins [17]. In *Xenopus laevis*, these present mostly N-linked, neutral glycans [52]. Reduction of PAS-positivity after β-elimination suggests that in *B. pachypus* O-linked glycans are also present, even if it is not possible to state whether they are attached to glycoproteins of the ZP family. An O-linked, galactosaminylated residual is also found in a *Xenopus laevis* ZP [17].

Similar to *B. pachypus* J₁, a strongly acidic layer adjacent to the FE is found in a number of species [4, 12, 53–55] and contains both carboxylated and sulfated glycoconjugates. Carboxylated glycans probably include sialic acid, as suggested by positivity to WGA lectin, which is involved in several functions, such as viscosity of the layer, three-dimensional conformational stability, protection from attack and degradation, hydration, osmotic regulation as well as sperm interaction [56–58]. Sulfated glycans can have functions similar to the sialylated (and, possibly, be present in the same chains [59]), and are known to inhibit polyspermy, whereas neutral glycans select sperms for penetration [8]. The presence of N-linked glycans suggests a structural role in keeping the three-dimensional organization of the layer and, possibly, the correct orientation of regulatory mole-

cules [59]. This is supported by the presence of fucosylated residues, suggesting the existence of a fucan fibrous superstructure to which globular glycoproteins are bound [5, 60]. Fucosylated residues can also act in reducing bacterial motility [61]. Mannosylated residues can be present in the core of the N-linked glycans [59]. The corresponding layer of *Xenopus laevis* resembles that of *B. pachypus* in having a fibrous structure and binding sites for WGA and ConA [5, 13, 17].

Further comparisons between our results and those of previous workers on other species are very difficult because of differences in the number of layers, techniques, interpretations, and terminology [3, 4].

In the J₂ layer of *B. pachypus*, the loose structure and the presence of mostly O-linked, acidic chains suggest a prevailing function in osmoregulation, hydration and spacing. Similar to J₁, WGA-binding could be explained by sialic acid and fucosylated residues could act against pathogens.

The J₃ layer is rich in proteoglycans, glycosaminoglycans and granules, probably made by globular glycoproteins. The highly acidic residues are probably involved in the same functions hypothesized for the previous layers; in particular, the relatively larger spaces among fibers can be important in terms of storing water and diffusible molecules, like sperm chemoattractants. Galactosaminyl and glucosaminyl residues found in J₃ layer can be explained by the presence of glycosaminoglycans [59].

The structure of the J₄ layer is similar to that of J₁, and carboxylated, mannosylated, galactosaminylated, glucosaminylated and fucosylated residues are also present. Thus, this layer could have functions similar to that of J₁. J₄ differs from J₁ in having galactosylated residues and in lacking sulfated and N-linked oligosaccharides. Galactose could be associated with N-acetylglucosamine in lactosamine structures that are common in O-linked glycan chains [62]. Polylactosamine chains, usually fucosylated or sialylated, can be involved in water retention, the presentation of glycan for interaction with lectin-like receptors, and in mucin/microorganism interactions [59].

J₅ is the outermost jelly coat and is probably involved in the adhesion and spacing between eggs, as well as in storing diffusible molecules involved in fertilization, something that has been observed in other species [14, 27, 63–65]. Its structural functions are suggested by the presence of N-linked glycans and mannosylated residues that can form the core of these glycans, as observed in the J₁ layer. Another important function of the J₅ layer is in interacting with the microorganisms coating the egg. It is unclear whether these organisms are epibionts, symbionts or patho-

gens. Oxygen supply and protection from excessive lighting functions have been proposed for several algae, some of which are even exclusive to amphibian egg jelly layers [3, 18], whereas a number of bacteria and molds feed on jelly layers and/or eggs and embryos [28]. Since microorganisms are found in *B. pachypus* J₅ layer, but not in the inner ones, it is probable that the mucins of the inner layers act as a barrier against them, as previously observed.

The previous functional interpretations are to be regarded as a preliminary attempt to link glycopatterns and structures; the functions proposed for a given layer can be shared with other ones. For example, fucosylated residues are present in all layers, so that structural and antibacterial functions could be suggested for each of them. Furthermore, functions of the same residues can differ between layers: acidic glycans can play a more important role in sperm interactions in the outer layers [8], whereas they could have a prevailing function in hydrosaline and acidic homeostasis in the inner ones.

B. pachypus presents a relatively high number of jelly layers in respect of other anurans [1, 3]. The number of layers itself can be linked to an increased protection against pathogens, since species with thicker jelly-coated eggs are less exposed to water mold infection than species with thinner coats [22]. Nonetheless, the jelly coat is not a sufficient barrier against penetration by herbicides like isoproturon [24], or insecticides like α -cypermethrin and endosulfan [25, 66], although it seems to reduce penetration by polycyclic aromatic hydrocarbons [21] and 2,4-D butoxyethyl ester [23], even though it is supposed that eggs of species with different jelly-coat thickness should differ in resistance to chemical penetration [21].

In conclusion, our study confirms the complex organization of the extracellular matrix of the amphibian egg and its species-specificity that can be fully understood only by integrating data from multiple approaches. Further studies are needed to fully understand the role of glycans in functions such as protection from pathogens and possible hosting for oxygen-supplying algae, as well as the effects of pollutants on their structure, and thus possible alterations of their protective functions.

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